

RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/536,880A
Source: IFWP
Date Processed by STIC: 8/14/06

ENTERED



IFWP

RAW SEQUENCE LISTING

DATE: 08/14/2006

PATENT APPLICATION: US/10/536,880A

TIME: 13:33:58

Input Set : A:\272924US0PCT.ST25.txt

Output Set: N:\CRF4\08142006\J536880A.raw

3 <110> APPLICANT: Onoue, Satomi
 4 Endo, Kousuke
 5 Matsumoto, Asami
 7 <120> TITLE OF INVENTION: PEPTIDES AND MEDICINAL COMPOSITIONS CONTAINING
 THE SAME
 9 <130> FILE REFERENCE: 272924US0PCT
 11 <140> CURRENT APPLICATION NUMBER: 10/536,880A
 12 <141> CURRENT FILING DATE: 2005-05-27
 14 <150> PRIOR APPLICATION NUMBER: PCT/JP03/14924
 15 <151> PRIOR FILING DATE: 2003-11-21
 17 <150> PRIOR APPLICATION NUMBER: JP 2002-344523
 18 <151> PRIOR FILING DATE: 2002-11-27
 20 <160> NUMBER OF SEQ ID NOS: 34
 22 <170> SOFTWARE: PatentIn version 3.3
 24 <210> SEQ ID NO: 1
 25 <211> LENGTH: 27
 26 <212> TYPE: PRT
 27 <213> ORGANISM: Artificial Sequence
 29 <220> FEATURE:
 30 <223> OTHER INFORMATION: Synthetic Peptide
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 34 <221> NAME/KEY: MISC_FEATURE
 35 <222> LOCATION: (5)..(5)
 36 <223> OTHER INFORMATION: Xaa = Val or Ile
 38 <220> FEATURE:
 39 <221> NAME/KEY: MISC_FEATURE
 40 <222> LOCATION: (8)..(8)
 41 <223> OTHER INFORMATION: Xaa = Asp, Glu or Ala
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 45 <222> LOCATION: (9)..(9)
 46 <223> OTHER INFORMATION: Xaa = Asn or Ser
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 49 <221> NAME/KEY: MISC_FEATURE
 50 <222> LOCATION: (11)..(11)
 51 <223> OTHER INFORMATION: Xaa = Thr or Ser
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 54 <221> NAME/KEY: MISC_FEATURE
 55 <222> LOCATION: (13)..(13)
 56 <223> OTHER INFORMATION: Xaa = Leu or Tyr
 58 <220> FEATURE:
 59 <221> NAME/KEY: MISC_FEATURE
 60 <222> LOCATION: (15)..(15)
 61 <223> OTHER INFORMATION: Xaa = Lys or Arg

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63 <220> FEATURE:
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65 <222> LOCATION: (17)..(17)
66 <223> OTHER INFORMATION: Xaa = Leu or Nle
68 <220> FEATURE:
69 <221> NAME/KEY: MISC_FEATURE
70 <222> LOCATION: (20)..(20)
71 <223> OTHER INFORMATION: Xaa = Lys or Arg
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74 <221> NAME/KEY: MISC_FEATURE
75 <222> LOCATION: (21)..(21)
76 <223> OTHER INFORMATION: Xaa = Lys or Arg
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79 <221> NAME/KEY: MISC_FEATURE
80 <222> LOCATION: (26)..(26)
81 <223> OTHER INFORMATION: Xaa = Ile or Val
83 <220> FEATURE:
84 <221> NAME/KEY: misc_feature
85 <222> LOCATION: (27)..(27)
86 <223> OTHER INFORMATION: Xaa can be any naturally occurring amino acid
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      91 1          5          10          15
      94 Xaa Ala Val Xaa Xaa Tyr Leu Ala Ala Xaa Xaa
      95          20          25
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108 <221> NAME/KEY: MOD_RES
109 <222> LOCATION: (28)..(28)
110 <223> OTHER INFORMATION: Amidation
112 <400> SEQUENCE: 2
114 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
115 1          5          10          15
118 Met Ala Val Lys Lys Tyr Leu Asn Ser Ile Leu Asn
119          20          25
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128 <223> OTHER INFORMATION: Synthetic Peptide
131 <220> FEATURE:
132 <221> NAME/KEY: MOD_RES
133 <222> LOCATION: (27)..(27)

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134 <223> OTHER INFORMATION: Amidation
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138 His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Lys Gln
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143          20          25
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157 <222> LOCATION: (38)..(38)
158 <223> OTHER INFORMATION: Amidation
160 <400> SEQUENCE: 4
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163 1          5          10          15
166 Met Ala Val Lys Lys Tyr Leu Ala Ala Val Leu Gly Lys Arg Tyr Lys
167          20          25          30
170 Gln Arg Val Lys Asn Lys
171          35
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186 <223> OTHER INFORMATION: Amidation
188 <400> SEQUENCE: 5
190 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Arg Gln
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194 Leu Ala Val Arg Arg Tyr Leu Asn Ser Ile Leu Asn Gly Arg Arg
195          20          25          30
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199 <211> LENGTH: 30
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201 <213> ORGANISM: Artificial Sequence
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204 <223> OTHER INFORMATION: Synthetic Peptide
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208 <221> NAME/KEY: MOD_RES
209 <222> LOCATION: (30)..(30)
210 <223> OTHER INFORMATION: Amidation
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214 His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Arg Gln
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218 Leu Ala Val Arg Arg Tyr Leu Ala Ala Val Leu Gly Arg Arg
219          20          25          30
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224 <212> TYPE: PRT
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233 <222> LOCATION: (38)..(38)
234 <223> OTHER INFORMATION: Amidation
236 <400> SEQUENCE: 7
238 His Ser Asp Gly Ile Phe Thr Asp Ser Tyr Ser Arg Tyr Arg Arg Gln
239 1          5          10          15
242 Leu Ala Val Arg Arg Tyr Leu Ala Ala Val Leu Gly Arg Arg Tyr Arg
243          20          25          30
246 Gln Arg Val Arg Asn Arg
247          35
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251 <211> LENGTH: 25
252 <212> TYPE: PRT
253 <213> ORGANISM: Artificial Sequence
255 <220> FEATURE:
256 <223> OTHER INFORMATION: Synthetic Peptide
259 <220> FEATURE:
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261 <222> LOCATION: (25)..(25)
262 <223> OTHER INFORMATION: Amidation
264 <400> SEQUENCE: 8
266 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
267 1          5          10          15
270 Met Ala Val Lys Lys Tyr Leu Asn Ser
271          20          25
274 <210> SEQ ID NO: 9
275 <211> LENGTH: 25
276 <212> TYPE: PRT
277 <213> ORGANISM: Artificial Sequence
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284 <221> NAME/KEY: MOD_RES
285 <222> LOCATION: (17)..(17)
286 <223> OTHER INFORMATION: Oxidation
288 <400> SEQUENCE: 9
290 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
291 1          5          10          15

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294 Met Ala Val Lys Lys Tyr Leu Asn Ser
295                20                25
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304 <223> OTHER INFORMATION: Synthetic Peptide
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309 <222> LOCATION: (23)..(23)
310 <223> OTHER INFORMATION: Amidation
312 <400> SEQUENCE: 10
314 His Ser Asp Ala Val Phe Thr Asp Asn Tyr Thr Arg Leu Arg Lys Gln
315 1                5                10                15
318 Leu Ala Val Lys Lys Tyr Leu
319                20
322 <210> SEQ ID NO: 11
323 <211> LENGTH: 23
324 <212> TYPE: PRT
325 <213> ORGANISM: Artificial Sequence
327 <220> FEATURE:
328 <223> OTHER INFORMATION: Synthetic Peptide
331 <220> FEATURE:
332 <221> NAME/KEY: MOD_RES
333 <222> LOCATION: (23)..(23)
334 <223> OTHER INFORMATION: Amidation
336 <400> SEQUENCE: 11
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339 1                5                10                15
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343                20
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355 <220> FEATURE:
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357 <222> LOCATION: (28)..(28)
358 <223> OTHER INFORMATION: Amidation
360 <400> SEQUENCE: 12
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363 1                5                10                15
366 Leu Ala Val Lys Lys Tyr Leu Ala Ala Ile Leu Asn
367                20                25
370 <210> SEQ ID NO: 13
371 <211> LENGTH: 30

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/536,880A

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220>

to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 5, 8, 9, 11, 13, 15, 17, 20, 21, 26, 27

Seq#:17; Xaa Pos. 17

VERIFICATION SUMMARY

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Input Set : A:\272924US0PCT.ST25.txt

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L:90 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

M:341 Repeated in SeqNo=1

L:495 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17 after pos.:16